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Smoothing for Discrete-Valued Time Series

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Abstract

We deal with smoothed estimators for conditional probability functions of discrete-valued time series $\{Y_t\}$ under two different settings. When the conditional distribution of Y_t given its lagged values falls in a parametric family and depends on exogenous random variables, a smoothed maximum (partial) likelihood estimator for the unknown parameter is proposed. While there is no prior information on the distribution, various nonparametric estimation methods have been compared and the adjusted Nadaraya-Watson estimator stands out as it shares the advantages of both Nadaraya-Watson and local linear regression estimators. The asymptotic normality of the proposed estimators has been established in the manner of sparse asymptotics, which shows that the proposed smoothed methods outperform their conventional, unsmoothed, parametric counterparts under very mild conditions. Simulation results lend further support to the above assertion. Finally, the new method is illustrated via a real data set concerning the relationship between the number of daily hospital admissions and the levels of pollutants in Hong Kong in 1994 – 1995. An ad hoc model selection procedure based on local AIC is proposed to select the significant pollutant indices.

Keywords: α -mixing; Adjusted Nadaraya-Watson estimator; Discrete-valued time series; Local AIC; Local linear smoother; Local partial likelihood; Nonparametric estimation; Smoothed maximum likelihood estimation; Sparse asymptotics.

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1 Introduction

Let $\{Y_t\}$ be a strictly stationary discrete-valued time series. We apply smoothing techniques to estimate the conditional probability function of Y_t given its lagged values in both parametric and nonparametric settings. The methods proposed are applicable when Y_t is either quantitative or ordinal categorical. *Such a variable can arise as a discretization of an underlying continuous variable or as an inherently discrete, but ordered, set of categories* (Simonoff, 1996, Section 6.1). In the former case, discrete values of Y_t have real physical meanings. If Y_t indeed represents a discretization of a continuous variable with smooth density function, the probability of $Y_t = i$ will be close to that of $Y_t = i + \Delta$ for small integer Δ . In the latter, the different values stand for different categories which have a natural ordering (for example, *very bad, bad, neutral, good, and very good*). An observation falling in one particular cell provides information about the probability of falling in its neighbors. Therefore, smoothing makes sense since we may assume that the probability function is ‘continuous’ (see the conditions (A3) and (B2) below) in both cases. The improvement using smoothing is most evident when the distribution is *sparse* in the sense that probability of Y_t falling each cell is small. To highlight this phenomenon, we develop an asymptotic approximation under the *sparse asymptotics* framework which assumes that the maximum value of the probability function converges to 0 when the sample size goes to infinity; see Simonoff (1985), Hall and Titterington (1987) and Simonoff (1996, Section 6.2). We will show that the proposed smoothed estimators have smaller asymptotic mean squared errors than those of conventional (non-smoothed) parametric estimators when the underlying distribution is sparse; see Remarks 2, 5 and 7 below.

In Section 2, we assume that the conditional distribution of Y_t given its past falls in a parametric family with the parameter depending on the value of Y_{t-1} and also an ‘exogenous’ variable. By assuming that the parameter is ‘continuous’ in Y_{t-1} (see the condition (A3) and Remark 1(iii) below), we estimate the parameter by maximizing a local (*i.e.* smoothed) partial likelihood function. We propose a simple and intuitively appealing bootstrap method to choose the bandwidth. The asymptotic normality of the estimator is established. In Section 3, various nonparametric kernel estimation methods for the conditional probability function of Y_t given Y_{t-1} are discussed. We are in favour of the adjusted Nadaraya-Watson estimator (Hall and Presnell, 1999; Hall, Wolff, and Yao, 1999) since it enjoys the same first-order asymptotic properties as

the local linear estimator and is always a proper probability function itself. The nonparametric setting in Section 3 is similar to the local polynomial estimation of continuous conditional density functions considered by Fan, Yao, and Tong (1996). However, the asymptotic theory is different since we estimate a probability function which converges to 0 itself. In Section 4, the proposed methods are illustrated through two simulated examples and a Hong Kong air pollution/disease data set.

Although it appears to us that smoothing techniques have not been used in analyzing discrete-valued time series data before, there has been a substantial amount of literature on their application to discrete data analysis. Aitchison and Aitken (1976), Bowman (1980), Titterton (1980) and Hall (1981) appear to be among the earliest. Lucid reviews on research in this direction can be found in Simonoff (1995; 1996, Chapter 6). The latest developments include Dong and Simonoff (1994) on boundary-corrected kernel estimation for sparse multinomial distributions, Aerts, Augustyns, and Janssen (1997) and Simonoff (1998) on local polynomial estimation of multinomial tables, and Faddy and Jones (1998) on semiparametric smoothing for discrete probability functions.

2 Smoothed Parametric Estimation

2.1 Maximum local-partial-likelihood estimator

Suppose that discrete-valued time series Y_t is influenced by an exogenous variable \mathbf{X}_t , $\{(\mathbf{X}_t, Y_t)\}$ forms a strictly stationary process, and Y_t takes non-negative integer values. We assume that the conditional probability of $Y_t = j$ given $(\mathbf{X}_t, \mathbf{X}_{t-1}, \dots, Y_{t-1} = i, Y_{t-2}, \dots)$ is $p(j; \mathbf{X}_t, \beta_i)$ which depends on $(\mathbf{X}_t, Y_{t-1} = i)$ only, where $p(\cdot; \cdot, \cdot)$ is of a given form and β_i is a unknown parameter vector. For example, the conditional distribution could be Poisson with mean $\mu(\mathbf{X}_t^T \beta_i)$, where $\mu(\cdot)$ is a known link function. Given observations $\{(\mathbf{X}_t, Y_t), 1 \leq t \leq n\}$, the log conditional likelihood function given \mathbf{X}_1 and Y_1 is

$$\sum_{t=2}^n \log\{p(Y_t; \mathbf{X}_t, \beta_{Y_{t-1}})\} + \sum_{t=2}^n \log\{f(\mathbf{X}_t; \mathbf{X}_{t-1}, \dots, \mathbf{X}_1, Y_{t-1}, \dots, Y_1)\},$$

where $f(X; Z)$ denotes the conditional probability density of X given Z . By maximizing the first sum in the above expression, we obtain the maximum (partial) likelihood estimators for

β_1, β_2, \dots . Such an estimator for β_i is in fact derived by maximizing

$$\sum_{t=2}^n \log\{p(Y_t; \mathbf{X}_t, \beta_i)\} I(Y_{t-1} = i), \quad (2.1)$$

which depends on the pairs (Y_{t-1}, Y_t) with $Y_{t-1} = i$ only. To make more efficient use of the available data, we define a smoothed (partial) likelihood function of β_i by replacing the indicator function $I(Y_{t-1} = i)$ by a kernel function:

$$\sum_{t=2}^n \log\{p(Y_t; \mathbf{X}_t, \beta_i)\} K_{n,h}(Y_{t-1} - i). \quad (2.2)$$

Maximizing the above smoothed likelihood, we obtain a smoothed estimator. In the above expression, $K(\cdot)$ is kernel function, $K_{n,h}(x) = h^{-1}K(\delta_n x/h)$, $h > 0$ is a bandwidth which controls the amount of smoothing used in estimation, and $\delta_n > 0$ reflects the sparseness of the underlying distribution; see, the condition (A3) and Remarks 1(ii) and (iii) in Section 2.3 below.

In view of the more attractive asymptotic properties of the local linear smoother relative to the local constant version (Fan, Farmen, and Gijbels, 1998), we propose to use the local linear estimator $\hat{\beta}_i$, where $(\hat{\beta}_i, \hat{\mathbf{a}})$ is the maximizer of

$$\ell_n = \sum_{t=2}^n \log\{p(Y_t; \mathbf{X}_t, \beta_i + \delta_n \mathbf{a}(Y_{t-1} - i))\} K_{n,h}(Y_{t-1} - i). \quad (2.3)$$

Obviously, the above approach can be applied to the case when the conditional probability of Y_t given its past is of the form $p(Y_t; \boldsymbol{\theta}_{Y_{t-1}})$ with known function p and unknown parameter $\boldsymbol{\theta}$. Further, the smoothing can also be incorporated into a quasilielihood approach of Wedderburn (1974) in an obvious manner. In fact the proposed smoothed (partial) likelihood function (2.2), although derived under a specified time series context, is of the form of the *local likelihood functions* explored by, among others, Tibshirani and Hastie (1987) and Fan, Farmen, and Gijbels (1998) for independent observations.

2.2 Bandwidth selection

The bandwidth h plays an important role in smoothing estimation. Most existing bandwidth selection methods were originally designed for continuous independent data, although some of them can be adapted to handle dependence in time series. For the problem discussed in this section, there is no natural way to derive an analogue to the cross validation method or its variations. Instead, we propose a simple bootstrap approach to choose h , which is easy to

implement, and takes into account of the dependence of the data in resampling. The bandwidth selected may be variable in the sense that different bandwidths may be used to estimate different β'_i s. The method is similar in spirit to those used by Hall, Wolff, and Yao (1999) and Polonik and Yao (2000) for estimation of (continuous) conditional distribution functions and conditional minimum volume sets.

We draw bootstrap samples conditionally on the given data $\{\mathbf{X}_t\}$ as follows. Let $\tilde{\beta}_i$ be the parametric estimator obtained from maximizing (2.1) for all i . (Some initial moving average smoothing may be applied to $\{\tilde{\beta}_i\}$ in the case that some cells contain few observations. Alternatively, $\tilde{\beta}_i$ can be obtained nonparametrically by maximizing (2.2) with a small bandwidth such that the biases are small.) Draw Y_0^* from the empirical (marginal) probability function of $\{Y_t\}$. For $t = 1, \dots, n$, draw Y_t^* from the (discrete) probability function $p(\cdot; \mathbf{X}_t, \tilde{\beta}_{Y_{t-1}^*})$. Define $\hat{\beta}_i^* \equiv \hat{\beta}_i^*(h)$ in the same way as $\hat{\beta}_i$ with $\{(\mathbf{X}_t, Y_t)\}$ replaced by $\{(\mathbf{X}_t, Y_t^*)\}$. To estimate β_i for a particular i , we choose h which minimizes the conditional expectation

$$E[|\hat{\beta}_i^*(h) - \tilde{\beta}_i| | \{(\mathbf{X}_t, Y_t)\}].$$

To speed up the computation, we may use one single bandwidth for the estimation of all of the β_i , and this single bandwidth minimizes

$$M(h) = \sum_i \hat{\pi}_i E[|\hat{\beta}_i^*(h) - \tilde{\beta}_i| | \{(\mathbf{X}_t, Y_t)\}], \quad (2.4)$$

where $\hat{\pi}_i$ is the relative frequency estimate for the marginal probability $P(Y_t = i)$. Some initial moving average may be applied in case some cells contain no observations. For example, we may replace $\hat{\pi}_i$ by the moving average of its three nearest neighbors (including itself) with the weights 1/2, 1/4 and 1/4.

2.3 Theoretical properties

We write $l(y; \mathbf{x}, \beta) = \log\{p(y; \mathbf{x}, \beta)\}$, and let

$$\dot{l}(y; \mathbf{x}, \beta) = \frac{\partial l(y; \mathbf{x}, \beta)}{\partial \beta}, \quad \text{and} \quad \ddot{l}(y; \mathbf{x}, \beta) = \frac{\partial^2 l(y; \mathbf{x}, \beta)}{\partial \beta \partial \beta^T}.$$

Define $\mu_j = \int u^j K(u) du$ and $\nu_j = \int u^j K(u)^2 du$. For matrix $\mathbf{A} = (a_{ij})$, $\|\mathbf{A}\| = (\sum a_{ij}^2)^{1/2}$. We use C to denote a finite positive constant which may be different at different places. We state some regularity conditions first.

(A1) The parameter β_i is identifiable in the sense that $p(\cdot; \mathbf{X}_t, \mathbf{z}_1) \neq p(\cdot; \mathbf{X}_t, \mathbf{z}_2)$ for any $\mathbf{z}_1 \neq \mathbf{z}_2$, and

$$\frac{\partial^2}{\partial \beta \partial \beta^T} E\{p(Y_t; \mathbf{X}_t, \beta_i) | \mathbf{X}_t, Y_{t-1} = i\} = E\left\{\frac{\partial^2}{\partial \beta \partial \beta^T} p(Y_t; \mathbf{X}_t, \beta_i) | \mathbf{X}_t, Y_{t-1} = i\right\}.$$

Further, all the third partial derivatives of $p(Y_t; \mathbf{X}_t, \beta_i)$ with respect to β_i are bounded by a random variable, say, $M(Y_t, \mathbf{X}_t)$, and $E\{M(Y_t, \mathbf{X}_t) | \mathbf{X}_t, Y_{t-1} = i\}$ is finite.

(A2) $\Sigma_i \equiv -E[E\{\ddot{l}(Y_t; \mathbf{X}_t, \beta_{Y_{t-1}}) | \mathbf{X}_t, Y_{t-1} = i\} | Y_{t-1} = i]$ is a positive definite matrix.

Further for some $\gamma > 2$,

$$E\left\{\|\dot{l}(Y_t; \mathbf{X}_t, \beta_i)\|^\gamma + \|\ddot{l}(Y_t; \mathbf{X}_t, \beta_i)\|^\gamma\right\} < \infty,$$

and $\|\Sigma_i - \Sigma_j\| \leq C\delta_n |j - i|$.

(A3) For $i = 0, 1, \dots$, $\pi_i \equiv P(Y_t = i) = \delta_n \int_i^{i+1} g(\delta_n x) dx$, where $g(\cdot)$ is a density function on $[0, \infty)$. Further, $\beta_i = \mathbf{b}(\delta_n i)$ and both $g(\cdot)$ and $\ddot{\mathbf{b}}(x) \equiv (\frac{\partial}{\partial x})^2 \mathbf{b}(x)$ are continuous in a neighborhood of $\delta_n i$.

(A4) The kernel function $K(\cdot)$ is bounded, symmetric and compactly supported.

(A5) The process $\{\mathbf{X}_t, Y_t\}$ is α -mixing with the mixing coefficient satisfying the condition $\alpha(k) = O(k^{-\beta})$, where $\beta > 2(\gamma - 1)/(\gamma - 2)$ for γ given in (A2) above.

(A6) As $n \rightarrow \infty$, $h \rightarrow 0$, $nh \rightarrow \infty$, and $\delta_n \rightarrow 0$.

Remark 1: Discussion of Conditions. (i) Both (A1) and (A2) are the standard conditions to ensure that the (unsmoothed) maximum likelihood estimator is consistent and asymptotically normal; see Lehmann and Casella (1998, Section 6.3). We need both of them for the consistency and asymptotic normality of $\hat{\beta}$ as well. Together with (A3) — (A6), they also ensure that the equation

$$\sum_{i=2}^n (1, \delta_n(Y_{t-1} - i))^T \otimes \dot{l}(Y_t; \mathbf{X}_t, \beta_i + \mathbf{a}\delta_n(Y_{t-1} - i)) K_{n,h}(Y_{t-1} - i) = 0 \quad (2.5)$$

has a solution which is a consistent estimator for β_i (see Theorem 1(i) below), where \otimes denotes the matrix Kronecker product. We refer $\hat{\beta}_i$ to such a solution hereafter.

(ii) Condition (A3) assumes $\pi_i = O(\delta_n)$. Note $\delta_n \rightarrow 0$ as $n \rightarrow \infty$. This reflects the fact that the *sparse asymptotics* depicts the performance of an estimator when the probability of Y_t falling into each cell is small. This is the case when the smoothing is most relevant.

(iii) The smooth condition imposed on both β_i and $g(\cdot)$ in (A3) reflects the fact that the proposed method is designed for the cases when the conditional probability function of Y_t given Y_{t-1} is ‘continuous’ in Y_{t-1} , which makes smoothing estimation relevant. The assumption that $\ddot{\mathbf{b}}(\cdot)$ is continuous ensures a nice asymptotic formula for the bias of $\widehat{\beta}_i$. If we replace it by

$$\|\beta_i - \beta_j\| \leq C\delta_n|i - j|,$$

$\widehat{\beta}_i$ is still asymptotically normal but with a bias of the order h .

(iv) The constant δ_n is introduced to reflect the sparseness of data. It does not add any extra complication in estimation of β'_i s. In fact in practical implementation we may fix δ_n at any value, the resulting discrepancy will be absorbed in the estimation of bandwidth h automatically; see (2.3).

(v) The requirement in (A4) that $K(\cdot)$ be compactly supported is imposed for the sake of brevity of proofs, and can be removed at the cost of lengthier arguments. In particular, the Gaussian kernel is allowed.

(vi) The α -mixing is one of the weakest mixing conditions for weakly dependent stochastic processes. Stationary time series or Markov chains fulfilling certain (mild) conditions are α -mixing with exponentially decaying coefficients; see Section 2.6.1 of Fan and Yao (2001). On the other hand, the assumption on the convergence rate of $\alpha(k)$ in (A5) is not the weakest possible and is imposed to simplify the proof.

Theorem 1. *Let conditions (A1)-(A6) hold. Suppose that $x_i = i\delta_n$ is bounded away from both 0 and ∞ as $n \rightarrow \infty$, and $g(x_i) > 0$. Then the following assertions hold.*

(i) *Equation (2.5) admits a root $\widehat{\beta}_i$ which converges to β_i in probability.*

(ii) *For any $\widehat{\beta}_i$ fulfilling (i) above,*

$$\widehat{\beta}_i - \beta_i = (nh)^{-1/2}\mathcal{N} + \frac{\mu_2 h^2}{2}\ddot{\mathbf{b}}(x_i) + o_P((nh)^{-1/2} + h^2),$$

where \mathcal{N} is a normal random vector with mean 0 and variance matrix $\nu_0 g(x_i)^{-1}\Sigma_i^{-1}$.

Remark 2: *Comparison with parametric estimator.* Under the condition (A3), $P(Y_t = i) = O(\delta_n)$. It can be shown that under this condition the parametric estimator $\widetilde{\beta}_i$ derived from maximizing (2.1) is asymptotically normal with mean 0 and variance of the order $1/(n\delta_n)$ instead of $1/n$, since the expected number of observations falling in each cell is of the order $n\delta_n$ instead of n . Theorem 1 shows that the asymptotic variance of the smoothed estimator $\widehat{\beta}_i$ is of the

order $1/(nh)$. Hence the asymptotic variance of $\widehat{\beta}_i$ converges to 0 *faster* than that of $\widetilde{\beta}_i$ for any bandwidth h for which $h/\delta_n \rightarrow \infty$. Theorem 1 indicates that the optimal bandwidth which minimizes the approximate mean squared error (AMSE) is of the order $n^{-1/5}$. (We define the AMSE as squared asymptotic bias plus asymptotic variance up to the first order.) With the optimal bandwidth, the AMSE of $\widehat{\beta}_i$ is of the order $n^{-4/5}$ which converges to 0 faster than the AMSE of $\widetilde{\beta}_i$ as long as $n\delta_n^5 \rightarrow 0$. This is a very mild condition. In the case that Y_t takes finite m values, $\delta_n = O(1/m)$. Therefore, the smoothed estimator $\widehat{\beta}_i$ will outperform parametric estimator $\widetilde{\beta}_i$ as long as $n = o(m^5)$.

Remark 3: *Asymptotic properties of conditional probabilities.* In the case that the exogenous variable \mathbf{X}_t is absent, Theorem 1 implies that

$$\begin{aligned} \frac{p(j, \widehat{\beta}_i)}{p(j, \beta_i)} - 1 &= p(j, \beta_i)^{-1} \dot{p}(j, \beta_i)^T (\widehat{\beta}_i - \beta_i) (1 + o_P(1)) \\ &= p(j, \beta_i)^{-1} \dot{p}(j, \beta_i)^T \left\{ (nh)^{-1/2} \mathcal{N} + \frac{\mu_2 h^2}{2} \ddot{\mathbf{b}}(x_i) + o_P((nh)^{-1/2} + h^2) \right\}, \end{aligned}$$

where $p(j, \beta_i) = P(Y_t = j | Y_{t-1} = i)$ and $\dot{p}(j, \beta_i) = \frac{\partial}{\partial \beta_i} p(j, \beta_i)$.

The proof of Theorem 1, as well as that of Theorem 2 in Section 3.3 below, is obtainable from the authors upon request.

3 Nonparametric Estimation

3.1 Estimators

We assume that data $\{Y_t, 1 \leq t \leq n\}$ are available from a strictly stationary discrete-valued time series, where Y_t takes integer values $\{1, \dots, m\}$ with $m < \infty$. Of interest is to estimate the conditional probability function

$$p_{ij} = P(Y_t = j | Y_{t-1} = i), \quad i, j = 1, \dots, m.$$

In fact the proposed methods below may also be extended to estimate higher-dimensional conditional probability function

$$p_{i_1, \dots, i_k, j} = P(Y_t = j | Y_{t-1} = i_1, \dots, Y_{t-k} = i_k)$$

for $k > 1$, which could be appealing when $\{Y_t\}$ is a k -th order Markov chain. However, such an extension is of limited practical value due to the difficulties associated with the curse of dimensionality.

Note that $p_{ij} = E\{I(Y_t = j) | Y_{t-1} = i\}$. This naturally leads to the Nadaraya-Watson (NW) estimator

$$\tilde{p}_{ij} = \frac{\sum_{t=2}^n I(Y_t = j) K_{mh}(Y_{t-1} - i)}{\sum_{t=2}^n K_{mh}(Y_{t-1} - i)}, \quad (3.1)$$

where $K(\cdot)$ is a kernel function, $K_{mh}(\cdot) = (mh)^{-1}K(\cdot/(mh))$, and $h > 0$ is a bandwidth which controls the amount of smoothness in estimation. The extreme case of $h = 0$ corresponds to the relative frequency estimate

$$\check{p}_{ij} = \frac{\sum_{t=2}^n I(Y_{t-1} = i, Y_t = j)}{\sum_{t=2}^n I(Y_{t-1} = i)} \quad \left(\frac{0}{0} \equiv 0 \right). \quad (3.2)$$

When $h > 0$, we use the information contained in the data (Y_{t-1}, Y_t) with $Y_t = j$ and Y_{t-1} close to i to estimate p_{ij} .

It is well known that an NW estimator exhibits boundary bias at both ends, which was addressed in Simonoff (1995) for categorical data, and it has more complicated asymptotic bias formula (see Remark 8 below). To attenuate these disadvantages, the obvious correction is to use the local linear estimator, defined as $\check{p}_{ij} = \hat{\alpha}$, due to its nice properties (Fan, 1993) such as mathematical efficiency, bias reduction and adaptation of edge effects, where $(\hat{\alpha}, \hat{\beta})$ minimizes

$$\sum_{t=2}^n [I(Y_t = j) - \alpha - \beta(Y_{t-1} - i)]^2 K_{mh}(Y_{t-1} - i). \quad (3.3)$$

However, \check{p}_{ij} is not constrained to lie between 0 and 1. In this aspect, the NW method is superior, since $\tilde{p}_{ij} \in [0, 1]$ and $\sum_j \tilde{p}_{ij} = 1$. We propose an ‘‘adjusted’’ version of the Nadaraya-Watson (ANW) estimator by combining the advantages from both NW and local linear estimators. The method was first introduced by Hall and Presnell (1999) for estimation of conditional mean functions and was used by Hall, Wolff, and Yao (1999) for the estimation of conditional distribution functions of continuous random variables.

The ANW approach is described as follows. Let $w_t(i)$, for $1 \leq t \leq n$, denote weights (function of the data Y_1, \dots, Y_{n-1} , as well as i) with the property that

$$w_t(i) \geq 0, \quad \sum_{t=2}^n w_t(i) = 1, \quad \text{and} \quad \sum_{t=2}^n w_t(i) (i - Y_{t-1}) K_{mh}(Y_{t-1} - i) = 0. \quad (3.4)$$

Of course, weights $w_t(i)$ satisfying these conditions are not uniquely defined, and we specify them concisely by maximizing $\prod_t w_t(i)$ subject to the constraints. As a result, $w_t(i)$ can be expressed as

$$w_t(i) = (n-1)^{-1} \{1 + \lambda (i - Y_{t-1}) K_{mh}(Y_{t-1} - i)\}^{-1},$$

where λ , a function of the data and i , is uniquely defined by (3.4). It is easily computed using a Newton-Raphson scheme. Then, the ANW estimator is defined by

$$\hat{p}_{ij} = \frac{\sum_{t=2}^n I(Y_t = j) w_t(i) K_{mh}(Y_{t-1} - i)}{\sum_{t=2}^n w_t(i) K_{mh}(Y_{t-1} - i)}. \quad (3.5)$$

Note particularly that $0 \leq \hat{p}_{ij} \leq 1$ and $\sum_j \hat{p}_{ij} = 1$. We show in Theorem 2 below that \hat{p}_{ij} is first-order equivalent to a local linear estimator which does not enjoy either of these properties.

3.2 Bandwidth selection

We may apply the generalized cross-validation (GCV) proposed by Wahba (1977) and Craven and Wahba (1979) to choose h . By ignoring the dependence on $\{I(Y_t = j)\}$ of the weight functions $\{w_i(t)\}$, it follows from (3.5) that

$$(\hat{p}_{Y_1, j}, \dots, \hat{p}_{Y_{n-1}, j})^\top = \mathbf{H} (I(Y_2 = j), \dots, I(Y_n = j))^\top,$$

where $\mathbf{H} = \mathbf{H}(h)$ is the $(n-1) \times (n-1)$ hat matrix. The GCV selects h which minimizes

$$\text{GCV}_j(h) = \left\{ 1 - \frac{\text{tr}(\mathbf{H})}{n} \right\}^{-2} \sum_{t=2}^n \left\{ I(Y_t = j) - \hat{p}_{Y_{t-1}, j} \right\}^2.$$

It is easy to see that $\text{tr}(\mathbf{H}) = K_{mh}(0) \sum_{l=2}^n [w_l(Y_{l-1}) / \sum_{t=2}^n w_t(Y_{l-1}) K_{mh}(Y_{t-1} - Y_{l-1})]$.

It is known that the GCV has a tendency of undersmoothing when $\text{tr}(\mathbf{H})/n$ is large, particularly for small sample size. To overcome this shortcoming, Hurvich, Simonoff, and Tsai (1998) proposed using the corrected version of Akaike information criterion (AICC)

$$\text{AICC}_j(h) = \log \left[\sum_{t=2}^n \left\{ I(Y_t = j) - \hat{p}_{Y_{t-1}, j} \right\}^2 \right] + \frac{2(\text{tr}(\mathbf{H}) + 1)}{(n-1) - \text{tr}(\mathbf{H}) - 2}.$$

It is easy to see that GCV and AICC are about the same when $\text{tr}(\mathbf{H})/n$ is small, which is typically the case in the context of analysing sparse discrete data.

Alternatively, the bootstrap approach described in Section 2.2 may also be adapted as follows. We generate a Markov chain $\{Y_t^*\}$ with the transition probability $\{\check{p}_{ij}\}$ defined in (2.1). Let $\hat{p}_{ij}^* \equiv \hat{p}_{ij}^*(h)$ be the estimator based on data $\{Y_t^*, 1 \leq t \leq n\}$, defined in the same manner as \hat{p}_{ij} . We use the bandwidth h in the estimator \hat{p}_{ij} , which minimizes the conditional expectation

$$E \left[\sum_{j=1}^m |\hat{p}_{ij}^*(h) - \check{p}_{ij}| \mid \{Y_t\} \right].$$

3.3 Theoretical properties

We impose the following regularity conditions. Write $x_i = i/m$ for $0 \leq i \leq m$.

(B1) As $n \rightarrow \infty$, $h \rightarrow 0$, $m \rightarrow \infty$, $nh/m \rightarrow \infty$, $m^2h^3 \rightarrow \infty$ and $nh^2 \rightarrow \infty$.

(B2) For $1 \leq i, j \leq m$,

$$\pi_{ij} \equiv P\{Y_{t-1} = i, Y_t = j\} = \frac{1}{m} \int_{x_{i-1}}^{x_i} \psi_j(x) dx, \quad (3.6)$$

where ψ_j is a positive function defined on $[0, 1]$ and has two continuous derivatives in a neighborhood of x_i . Further, the second derivative of the density function $\psi(x) \equiv m^{-1} \sum_{j=1}^m \psi_j(x)$ is bounded by a constant independent of m in the same neighborhood.

(B3) $K(\cdot)$ is a bounded, symmetric density function with a compact support.

(B4) The process $\{Y_t\}$ is strictly stationary and α -mixing with the mixing coefficient $\alpha(k) = O(k^{-\beta})$ as $k \rightarrow \infty$, where $\beta > 2$ is a constant.

Remark 4. *Discussion of Conditions.* Note that $p_{ij} = \pi_{ij}/\pi_i$, where $\pi_i = \int_{x_{i-1}}^{x_i} \psi(x) dx$. It is easy to see that the condition (B2) implies that $p_{ij} = O(m^{-1})$. Since we only deal with sparse distributions, we assume the number of categories m tends to ∞ as the sample size $n \rightarrow \infty$. We assume ψ_j (therefore also ψ) has two continuous derivatives in order to pursue good asymptotic properties. As long as ψ_j is Lipschitz continuous in a neighborhood of x_i , our estimators are still asymptotically normal but with larger biases (of the order of h).

We only present the asymptotic normality for the adjusted Nadaraya-Watson estimator \hat{p}_{ij} in the theorem below, and compare it with other methods in the discussion followed. Let $\varphi_j(x) = \psi_j(x)/\psi(x)$, and μ_j and ν_j be the same as in Section 2.3. We denote by $\dot{\varphi}(\cdot)$ and $\ddot{\varphi}(\cdot)$ the first two derivatives of $\varphi(\cdot)$ respectively.

Theorem 2. *Suppose that the conditions (B1) – (B4) hold, and $x_i = i/m$ is bounded away from both 0 and 1 as $n \rightarrow \infty$. Then for any $1 \leq j \leq m$,*

$$\hat{p}_{ij}/p_{ij} - 1 = \left(\frac{m\nu_0}{nh\psi_j(x_i)} \right)^{1/2} Z + h^2 \frac{\mu_2 \ddot{\varphi}_j(x_i)}{2\varphi_j(x_i)} + o_P(\{m/(nh)\}^{1/2} + h^2) + O_P(m^{-1}), \quad (3.7)$$

where Z stands for a standard normal random variable.

Remark 5: *Sparse asymptotics.* Since $p_{ij} \rightarrow 0$, we consider the asymptotic normality of \widehat{p}_{ij}/p_{ij} instead of p_{ij} . Note that the number of observations falling in each category is of the order n/m which can be viewed as the equivalent sample size in a usual asymptotic setting where $p_{ij} > 0$ is fixed (*i.e.* not converges to 0 as $n \rightarrow \infty$). This explains that the convergence rate in the above theorem is $(nh/m)^{1/2}$ instead of the conventional $(nh)^{1/2}$. Consequently, the optimal bandwidth h which minimizes the approximate mean squared error (AMSE) is of the order $(m/n)^{1/5}$.

Remark 6: *Comparison with relative frequency estimator.* If the conditions (B2) and (B4) holds and both n/m and m tend to infinity, it holds that $(\check{p}_{ij}/p_{ij} - 1)$ is asymptotically normal with mean 0 and asymptotic variance of the order m^2/n . Note that the asymptotic variance of \widehat{p}_{ij}/p_{ij} is of the order $m/(nh)$. Hence with any h such that $mh \rightarrow \infty$, the asymptotic variance of the smoothed estimator \widehat{p}_{ij} converges to 0 *faster* than that of the unsmoothed estimator \check{p}_{ij} . If we use the optimal bandwidth $h_{opt} \propto (m/n)^{1/5}$, for which $mh \rightarrow \infty$ under very mild restriction $n = o(m^6)$, the above assertion on the asymptotic variance also holds for the AMSE.

Remark 7: *Comparison with other nonparametric estimators.* The local linear estimator \check{p}_{ij} derived from (3.3) admits the same asymptotic expression as (3.7). For the NW estimator \widetilde{p}_{ij} defined in (3.1), the asymptotic expression still holds if we replace the second term on the right hand side of (3.7) (*i.e.* the bias) by

$$\frac{1}{2}h^2\mu_2[\ddot{\varphi}_j(x_i)/\varphi_j(x_i) + 2\dot{\varphi}_i(x_i)\dot{\psi}(x_i)/\{\psi(x_i)\varphi_j(x_i)\}],$$

which has one more term. Since all smoothed estimators \widehat{p}_{ij} , \check{p}_{ij} and \widetilde{p}_{ij} share the same asymptotic variance and the same order (*i.e.* $O(h^2)$) biases, the assertions on the superiority over unsmoothed estimator \check{p}_{ij} in Remark 6 above are also valid for \check{p}_{ij} and \widetilde{p}_{ij} .

Remark 8: *Boundary properties.* Let i be a boundary point, *i.e.* $i/m = ch$ for some $c \in (0, 1)$.

Then it can be proved that

$$\begin{aligned} \widehat{p}_{ij}/p_{ij} - 1 &= \left(\frac{m}{nh}\right)^{1/2} \frac{\eta_2(c)^{1/2}}{\eta_1(c)\{\varphi_j(0)\psi(0)\}^{1/2}} Z + h^2 \frac{\eta_0(c)\ddot{\varphi}_j(0)}{2\eta_1(c)\varphi_j(0)} \\ &\quad + o_P\left(\left(\frac{m}{nh}\right)^{1/2} + h^2\right) + O_P(m^{-1}), \\ \check{p}_{ij}/p_{ij} - 1 &= \left(\frac{m}{nh}\right)^{1/2} \frac{\{\int_{-c}^1 (\mu_{c,2} - \mu_{c,1}u)^2 K(u)^2 du\}^{1/2}}{(\mu_{c,0}\mu_{c,2} - \mu_{c,1}^2)\{\varphi_j(0)\psi(0)\}^{1/2}} Z \\ &\quad + h^2 \frac{\ddot{\varphi}_j(0)}{2\varphi_j(0)} \frac{\mu_{c,2}^2 - \mu_{c,1}\mu_{c,3}}{\mu_{c,0}\mu_{c,2} - \mu_{c,1}^2} + o_P\{\{m/(nh)\}^{1/2} + h^2\} + O_P(m^{-1}), \end{aligned}$$

$$\tilde{p}_{ij}/p_{ij} - 1 = \left(\frac{m}{nh}\right)^{1/2} \frac{\{\int_{-c}^1 K(u)^2 du\}^{1/2}}{\mu_{c,0}\{\varphi_j(0)\psi(0)\}^{1/2}} Z + h \frac{\mu_{c,1}\ddot{\varphi}_j(0)}{\mu_{c,0}\varphi_j(0)} + o_P\left(\left(\frac{m}{nh}\right)^{1/2} + h\right) + O_P(m^{-1}),$$

where $\mu_{c,k} = \int_{-c}^1 u^k K(u) du$, and

$$\eta_0(c) = \int_{-c}^1 \frac{u^2 K(u)}{1 - \lambda_c u K(u)} du, \quad \eta_k(c) = \int_{-c}^1 \frac{K(u)^k}{\{1 - \lambda_c u K(u)\}^k} du \quad (k = 1, 2).$$

In the above expressions, λ_c is the root of equation $\int_{-c}^1 u K(u) / \{1 - \lambda u K(u)\} du = 0$. These results confirm that both ANW estimator and local linear estimator are also boundary-adaptive in the sparse asymptotics in the sense that the asymptotic variances and biases are of the same orders as at the inner points. Therefore, both \hat{p}_{ij} and \check{p}_{ij} perform better than the unsmoothed estimator \check{p}_{ij} even at the boundary points; see Remark 6 above.

Remark 9: *Comparison with smoothed parametric estimators.* In the scenario described in Remark 3, we may also apply nonparametric estimation for $p_{ij} \equiv p(j, \beta_i)$ if Y_t takes finite values $1, \dots, m$. Theorem 2 shows that the bias of the resulting estimator is of the order h^2 , which is the same as the smoothed parametric estimator; see Remark 3. The asymptotic approximation for the variance is

$$V_1(j, i) \equiv \text{Var}\{\hat{p}_{ij}/p_{ij} - 1\} \approx \frac{m}{nh} \frac{\nu_0}{\psi_j(i/m)} = \frac{1}{nh} \frac{\nu_0}{m p_{ij} \pi_i} \{1 + o(1)\}.$$

The last equality follows from the relation that $\psi_j(i/m) \sim m^2 \pi_{ij} = m^2 p_{ij} \pi_i$. On the other hand, the smoothed estimation yields the asymptotic variance

$$\begin{aligned} V_2(j, i) &\equiv \text{Var}\left\{\frac{p(j, \hat{\beta}_i)}{p(j, \beta_i)} - 1\right\} \approx \frac{\nu_0}{nhg(\delta_n i) p^2(j, \beta_i)} \dot{p}(j, \beta_i)^T \Sigma_i^{-1} \dot{p}(j, \beta_i) \\ &= \frac{\nu_0 \delta_n}{nh \pi_i p^2(j, \beta_i)} \dot{p}(j, \beta_i)^T \Sigma_i^{-1} \dot{p}(j, \beta_i) \{1 + o(1)\}, \end{aligned}$$

see Remark 3 and Theorem 1. To facilitate the comparison between two estimators, we consider the average asymptotic approximations

$$E\{V_1(Y_t, i) | Y_{t-1} = i\} \approx \frac{\nu_0}{nh \pi_i},$$

and

$$E\{V_2(Y_t, i) | Y_{t-1} = i\} \approx \frac{\delta_n \nu}{nh \pi_i} \text{tr}[\Sigma_i^{-1} E\{\frac{\dot{p}(Y_t, \beta_i) \dot{p}(Y_t, \beta_i)^T}{p^2(Y_t, \beta_i)} | Y_{t-1} = i\}] = \frac{\nu_0}{nh \pi_i} \delta_n d,$$

where d is the dimensionality of β , which is a fixed constant. This shows that the ratio of the variance for the smoothed parametric estimator to its nonparametric counterpart converges to 0

at the rate δ_n . Therefore we should use the parametric approach whenever there are grounds to do so. Our empirical study also confirms the superior performance of the smoothed parametric estimation; see Example 2 in Section 4.

4 Numerical Properties

To assess the finite sample performance of the proposed smoothed estimators, we apply them to two simulated examples and one real data set. For the simulation models, we compare the smoothed estimators with their parametric counterparts. We also compare the nonparametric and parametric smoothed estimators in Example 2. Throughout this section, the Epanechnikov kernel $K(u) = 0.75(1 - u^2)I(|u| \leq 1)$ is used.

Example 1. First we consider a Poisson time series model constructed as follows. Let $\{X_t\}$ be a sequence of independent identically distributed random variables from uniform $[-1, 1]$. Given $\{(X_{s+1}, Y_s), s \leq t\}$, the conditional distribution of Y_{t+1} is Poisson with the mean $\lambda(X_{t+1}, Y_t)$, where $\lambda(x, i) = \exp(\beta_{1i} + \beta_{2i} x)$, and

$$\beta_{1i} = 4 - 2.49 \exp\left\{-\frac{(i - 25)^2}{512}\right\}, \quad \beta_{2i} = -1.5 \sin(\pi i/25).$$

Obviously, $\{(X_t, Y_t)\}$ is a homogeneous Markov chain. Figure 1a plots a sample of time series $\{Y_t\}$ of length 200. We repeat the simulation 400 times for each of the sample sizes $n = 100, 200$ and 400. For each realization, we calculate the smoothed estimator $(\hat{\beta}_{1i}, \hat{\beta}_{2i})$ which maximizes smoothed likelihood function defined as in (2.3), as well as the parametric estimator $(\tilde{\beta}_{1i}, \tilde{\beta}_{2i})$ which is derived by maximizing the likelihood function defined as in (2.1). For the smoothed estimator, we set $\delta_n = 0.1$ and use the bandwidth h which minimizes $M(h)$ defined in (2.4). Figure 1b presents boxplots of the mean absolute deviation errors (ADE)

$$\mathcal{E} = \sum_i \hat{\pi}_i \left\{ |\hat{\beta}_{1i} - \beta_{1i}| + |\hat{\beta}_{2i} - \beta_{2i}| \right\}, \quad (4.1)$$

where $\hat{\pi}_i$ is the relative frequency estimator for the marginal probability $P(Y_t = i)$. The measure \mathcal{E} decreases as the sample size increases. Figure 1c depicts boxplots of the mean *relative* absolute deviation errors

$$\mathcal{E}_r = \sum_i \hat{\pi}_i \frac{|\hat{\beta}_{1i} - \beta_{1i}| + |\hat{\beta}_{2i} - \beta_{2i}|}{|\tilde{\beta}_{1i} - \beta_{1i}| + |\tilde{\beta}_{2i} - \beta_{2i}|}. \quad (4.2)$$

Note that $\mathcal{E}_r < 1$ implies that the smoothed estimation outperforms its parametric counterpart. Figure 1c shows that the improvement of using the smoothed method is substantial, although

the difference decreases as n increases. The latter indicates that the smoothed method is more relevant for small sample sizes, although for sample size as large as 400 it is still significantly better than the unsmoothed method in this example. Figures 1d and 1e plot the typical example of estimated β_{1i} and β_{2i} against i , together with their true values. Typical example is selected such that the corresponding \mathcal{E} is equal to the median in the 400 replicated simulations. Figure 1f plots $M(h)$ defined in (2.4) versus bandwidth h , which indicates that the optimal bandwidth is 0.35 for this typical example.

Example 2. We consider a Markov chain time series $\{Y_t\}$ generated as follows. Given $\{Y_s, s \leq t\}$, the conditional distribution of Y_{t+1} is $\text{binomial}(m, p(Y_t))$, where

$$\text{logit}(p(i)) = i/m - (i/m)^2 \equiv \beta_i.$$

Figures 2a – 2c plot segments of time series $\{Y_t\}$ of length 200 for $m = 5, 10$ and 20 respectively. For each of the sample sizes $n = 100, 200,$ and 400 and $m = 5, 10,$ and 20, we repeat the simulation 400 times. For each realization, we compute the ANW estimator $\{\hat{p}_{ij}\}$ defined in (3.5) with the GCV bandwidth (see Section 3.2) and the relative frequency estimator $\{\check{p}_{ij}\}$ given in (3.2). Figure 2d presents boxplots of mean absolute deviation errors (ADE)

$$\mathcal{E} = \sum_i \hat{\pi}_i \sum_j |\hat{p}_{ij} - p_{ij}| \quad (4.3)$$

for $m = 5$ (the three panels on the left), 10 (the three panels in the middle) and 20 (the three panels on the right), where $\hat{\pi}_i$ is the relative frequency estimator for the marginal probability $P(Y_t = i)$. It is clear that \mathcal{E} decreases as the sample size n increases. Figure 2e displays the boxplots of the mean *relative* absolute deviation errors

$$\mathcal{E}_r = \sum_i \hat{\pi}_i \frac{\sum_j |\hat{p}_{ij} - p_{ij}|}{\sum_j |\check{p}_{ij} - p_{ij}|}. \quad (4.4)$$

It holds always that $\mathcal{E}_r < 1$ in Figure 2e. This indicates that the nonparametric estimator always performs better than the relative frequency estimator in this example, although the difference between the two methods decreases as the sample size n increases. Note that as m increases, \mathcal{E}_r decreases. This illustrates that the more sparse the distribution is, the more relevant the smoothing is. Figures 2g – 2l plot the typical example of estimated \hat{p}_{ij} (dotted line) and \check{p}_{ij} (dashed line) against j for $n = 200$ and $m = 10$, together with the true value p_{ij} (solid line), for five cases: from $i = 1$ to $i = 5$. They show clearly that the ANW estimates are more accurate

than the relative frequency estimates. Typical examples are selected such that the corresponding \mathcal{E} 's are equal to the medians in the 400 replicated simulations.

To illustrate the superior performance of the smoothed parametric estimation over the purely nonparametric estimation, we apply smoothed maximum likelihood method (see (2.3)) to obtain the smoothed parametric estimator $\widehat{\beta}_i$, and then compare directly the derived estimator

$$p_{ij}^* = \binom{m}{j} \left(\frac{e^{\widehat{\beta}_i}}{1 + e^{\widehat{\beta}_i}} \right)^j \left(1 - \frac{e^{\widehat{\beta}_i}}{1 + e^{\widehat{\beta}_i}} \right)^{m-j}$$

with the nonparametric estimator \widehat{p}_{ij} obtained above. Figure 2f presents the boxplots of values of

$$\mathcal{E}^* = \sum_i \widehat{\pi}_i \sum_j |p_{ij}^* - p_{ij}| \quad (4.5)$$

in the 400 replications. A direct comparison with Figure 2d indicates that that overall p_{ij}^* is much more accurate than \widehat{p}_{ij} . Furthermore, Figure 2g displays boxplots of the mean relative ADEs for p_{ij}^* over \widehat{p}_{ij}

$$\mathcal{E}_r^* = \sum_i \widehat{\pi}_i \frac{\sum_j |p_{ij}^* - p_{ij}|}{\sum_j |\widehat{p}_{ij} - p_{ij}|}, \quad (4.6)$$

which indicates a significant gain from using parametric model, as \mathcal{E}_r^* is always smaller than 1 and in fact is smaller than 0.4 in most cases. The finding here reinforces the theoretical results in Remark 9.

Example 3. Finally we apply the proposed methods to Hong Kong environmental data. The data were collected daily in Hong Kong from January 1, 1994 to December 31, 1995 (courtesy of Professor T. S. Lau). Of interest is to examine the relationship between the total number of daily hospital admissions (Y_t) for circulatory and respiratory problems and the levels of pollutants. Figure 3a displays the number of daily hospital admissions. The covariates are taken as the levels of pollutants sulphur dioxide X_{1t} (in $\mu g/m^3$), nitrogen dioxide X_{2t} (in $\mu g/m^3$) and dust X_{3t} (in $\mu g/m^3$). The correlation coefficient between X_{2t} and X_{3t} is 0.782, which is quite high.

Fan and Zhang (1999) used a varying-coefficient model

$$Y_t = a_1(t) + a_2(t) X_{1t} + a_3(t) X_{2t} + a_4(t) X_{3t} + \varepsilon_t \quad (4.7)$$

to fit the daily data. Cai, Fan and Li (2000) considered a Poisson regression model to the weekly data with mean $\lambda(t, \mathbf{X}_t)$ given by

$$\log\{\lambda(t, \mathbf{X}_t)\} = a_1(t) + a_2(t) X_{1t} + a_3(t) X_{2t} + a_4(t) X_{3t}. \quad (4.8)$$

While the fitted models are interesting, both approaches ignore the auto-dependence in the data; see Figure 3b. Cai, Fan and Li (2000) argued that the autocorrelation of the response variable is not strong for the weekly data. They also pointed out that X_{3t} in the above model is not significant according to a goodness-of-fit test.

Following the lead of Cai, Fan, and Li (2000), we model the daily hospital admissions with Poisson distributions. However, instead of letting the parameter in the link function vary with respect to the time, we assume it is a function of the number of patients in the immediate past. By modeling data in such a way, we are able to incorporate the dependence in the time series into the model. We assume that the number of daily admissions Y_t follows a Poisson distribution with the mean $\lambda_i(\mathbf{X}_t, \boldsymbol{\beta}_i)$, conditionally on its lagged values $Y_{t-1} = i, Y_{t-2}, Y_{t-3}, \dots$, and the levels of pollutants, where $\lambda_i(\cdot, \cdot)$ is given as

$$\log\{\lambda_i(\mathbf{X}_t, \boldsymbol{\beta}_i)\} = \beta_{1i} + \beta_{2i} X_{1t} + \beta_{3i} X_{2t} + \beta_{4i} X_{3t}. \quad (4.9)$$

This model differs from that of Cai, Fan, and Li (2000) since the parameters β 's now vary with respect to the immediately lagged value Y_{t-1} rather than time t . Note that in this model the dependence of Y_t on Y_{t-1} has been also reflected indirectly by its association with pollutants X_{jt} for $j = 1, 2$, and 3 . Therefore, there is a genuine need to delete the insignificant variables in (4.9). To this end, we propose an ad hoc procedure based on the *local* AIC as follows. In general, the AIC is defined as

$$-2(\text{maximized log likelihood}) + 2(\text{number of estimated parameters}).$$

See Akaike (1973). From (2.2), we may define the local AIC at $Y_{t-1} = i$ for this example as

$$\text{AIC}_i(d) = 2 \sum_{t=2}^n \left[\lambda_i(\mathbf{X}_t, \hat{\boldsymbol{\beta}}_i) - Y_t \log \left\{ \lambda_i(\mathbf{X}_t, \hat{\boldsymbol{\beta}}_i) \right\} \right] K_{n,h}(Y_{t-1} - i) + 2d,$$

where d is the number of nonzero components of $\boldsymbol{\beta}_i$. By minimizing $\text{AIC}_i(d)$ over d , we derive an ‘optimum’ model for the conditional distribution of Y_t given $Y_{t-1} = i$. However, we are interested in the global form of mean in this example. We simply choose the model which minimizes the average local AIC defined as

$$\text{AIC}(d) = \sum_i \hat{\pi}_i \text{AIC}_i(d), \quad (4.10)$$

where $\hat{\pi}_i$ is the relative frequency estimate for $P(Y_t = i)$. By taking δ_n to be 0.05 and using the bandwidth selector described in Section 2.2, we computed the AIC values defined in (4.10) for

Table 1: The AIC values for the eight candidate models.

Model with covariate(s)	no X_j 's	X_1	X_2	X_3	(X_1, X_2)	(X_1, X_3)	(X_2, X_3)	(X_1, X_2, X_3)
AIC-min(AIC)	6.891	6.740	1.134	0.382	0.800	0.000	0.852	1.559

all eight possible models (with no interactions), which are reported in Table 1. This leads to the selected model

$$\log\{\lambda_i(\mathbf{X}_t, \boldsymbol{\beta})\} = \beta_{1i} + \beta_{2i} X_{1t} + \beta_{4i} X_{3t}. \quad (4.11)$$

The corresponding optimal bandwidth is 1.10. Figure 3c plots estimated intercept $\hat{\beta}_{1i}$ against i (the value of Y_{t-1}) and Figure 3d plots the estimated coefficients $\hat{\beta}_{2i}$ and $\hat{\beta}_{4i}$ against i . The fitted model for λ_i is dominated by the intercept $\hat{\beta}_{1i}$ which increases monotonically as i increases. This indicates clearly that the (conditional) distribution of Y_t depends on Y_{t-1} . Further, Y_t tends to be large when Y_{t-1} is large. This reflects the auto-dependence observed in Figure 3b.

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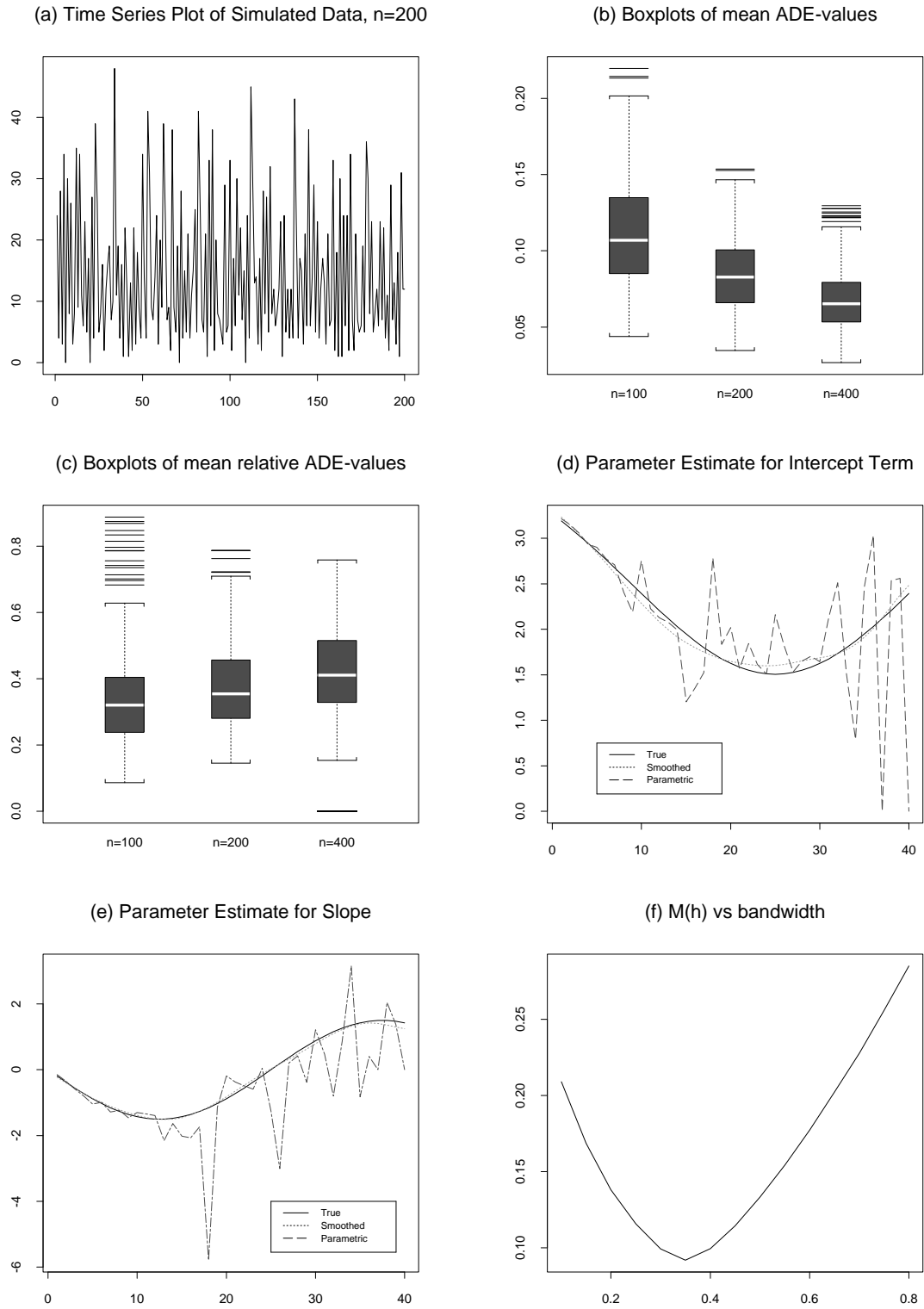
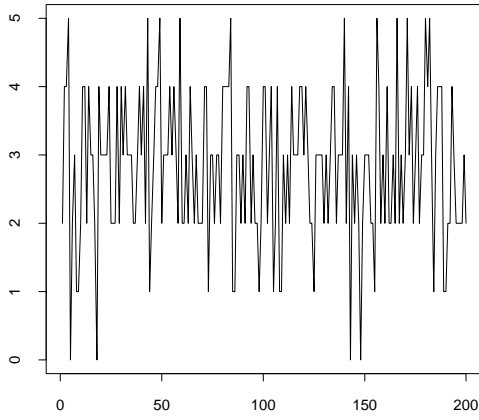
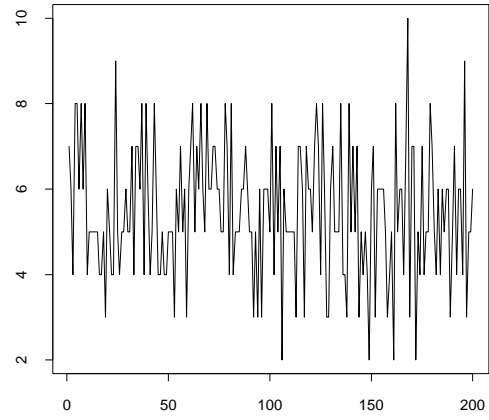


Figure 1: *Simulation results for Example 1. (a) Plot of a time series $\{Y_t\}$ of length 200. (b) Boxplot of 400 values of \mathcal{E} defined in (4.1). (c) Boxplot of 400 values of \mathcal{E}_r defined in (4.2). (d) Plot of true β_{1i} (solid line), $\hat{\beta}_{1i}$ (dot line) and $\tilde{\beta}_{1i}$ (dashed line) against i . (e) Plot of true β_{2i} (solid line), $\hat{\beta}_{2i}$ (dotted line) and $\tilde{\beta}_{2i}$ (dashed line) against i . (f) Plot of $M(h)$ against bandwidth h .*

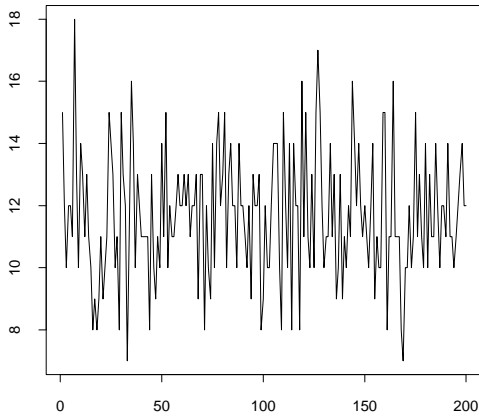
(a) Time Series Plot of Simulated Data, $n=200$, $m=5$



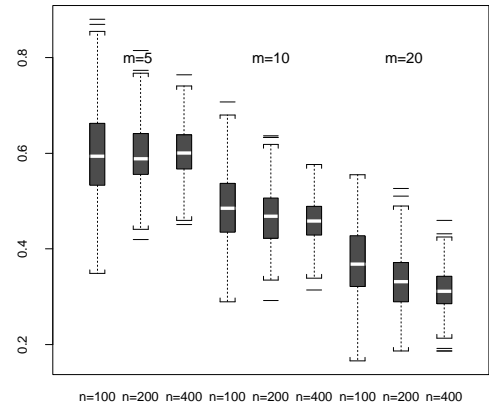
(b) Time Series Plot of Simulated Data, $n=200$, $m=10$



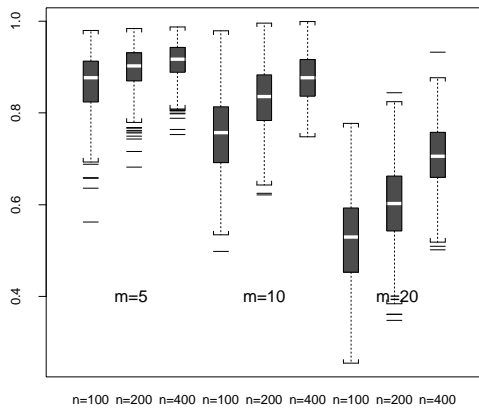
(c) Time Series Plot of Simulated Data, $n=200$, $m=20$



(d) Boxplots of mean ADE-values



(e) Boxplot of mean relative ADE-values



(f) Boxplots of mean ADE-values

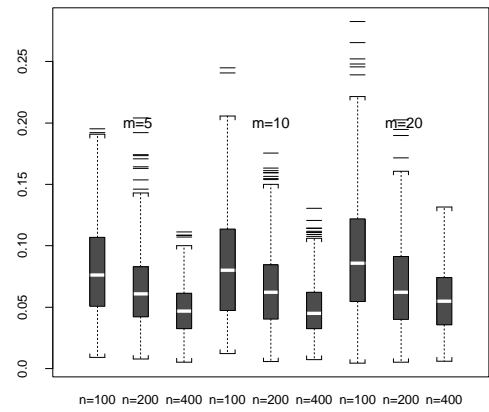


Figure 2: Simulation results for Example 2. (a)-(c) Plot of a time series $\{Y_t\}$ of length 200 for $m = 5, 10$ and 20 . (d) Boxplot of 400 values of \mathcal{E} defined in (4.3). (e) Boxplot of 400 values of \mathcal{E}_r defined in (4.4). (f) Boxplot of 400 values of \mathcal{E}^* defined in (4.5).

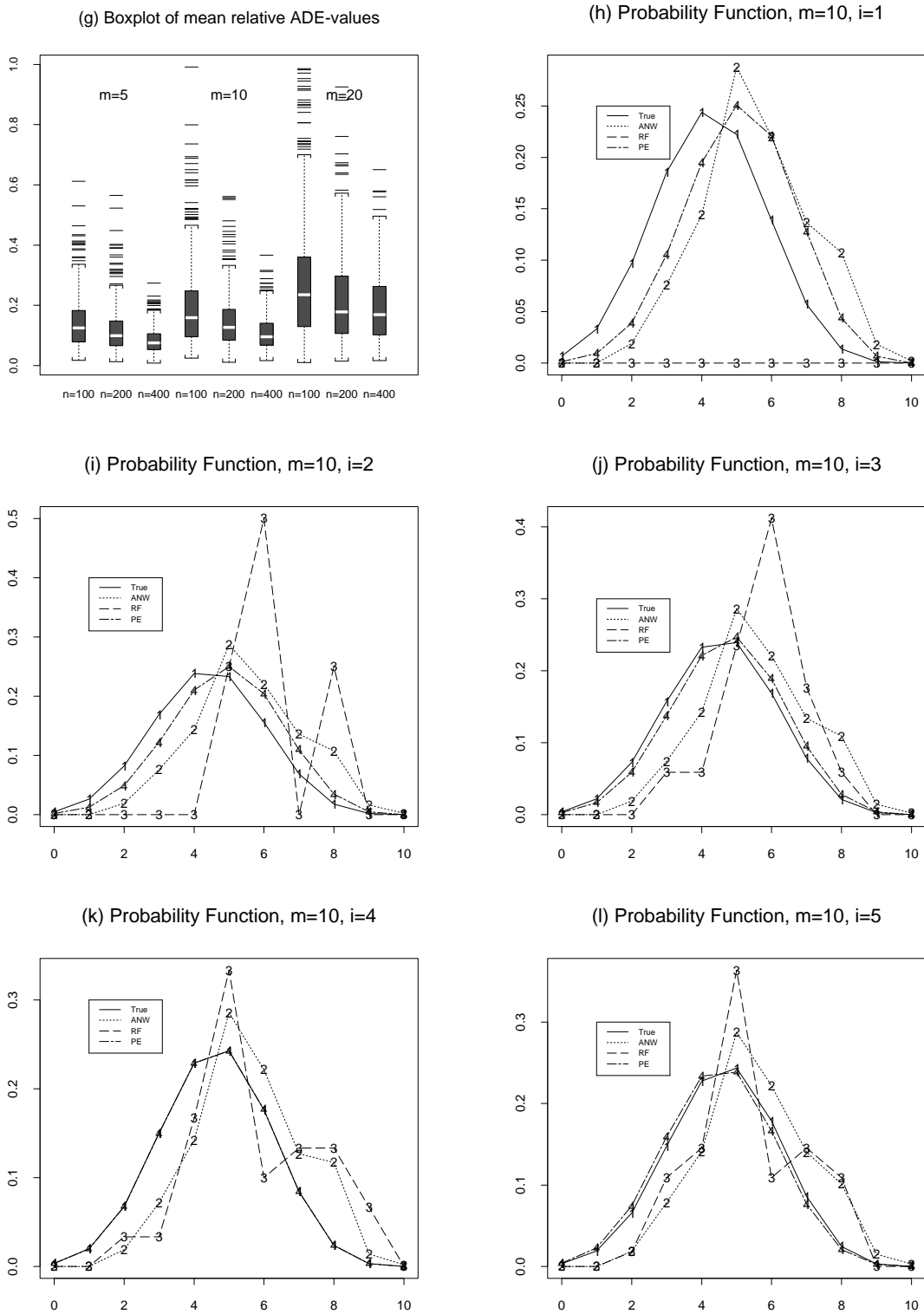
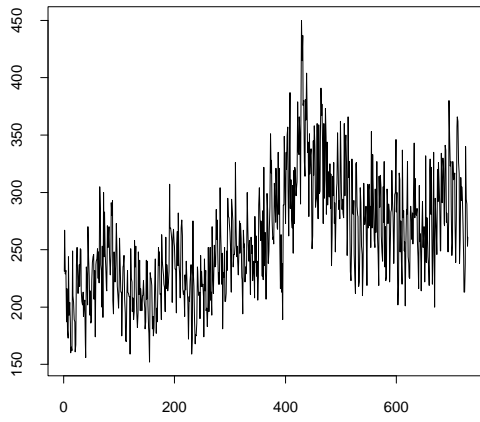
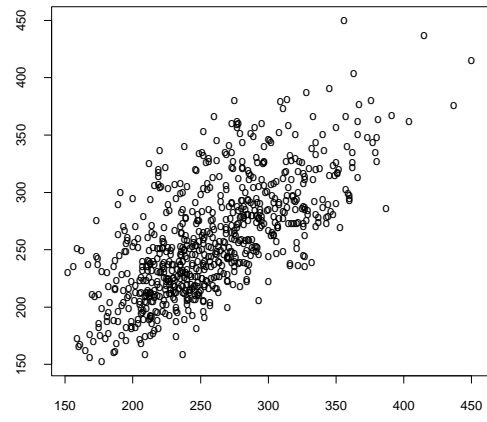


Figure 2: (Continued.) (g) Boxplot of 400 values of \mathcal{E}_r^* defined in (4.6). (h)-(l) Plot of true p_{ij} (solid line), \hat{p}_{ij} (dotted line), \check{p}_{ij} (dashed line), and \tilde{p}_{ij}^* (dotted-dashed line) against j for $n = 200$ $m = 10$. (h) $i = 1$, (i) $i = 2$, (j) $i = 3$, (k) $i = 4$, and (l) $i = 5$.

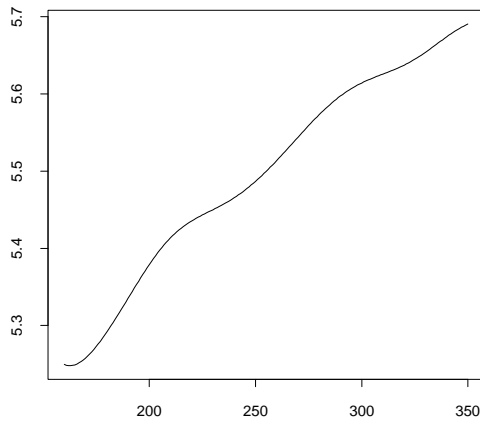
(a) Time Series Plot of Daily Hospital Admissions



(b) Scatterplot of Y_t vs Y_{t-1}



(c) Estimated Coefficient: β_1



(d) Estimated Coefficients: β_2 and β_4

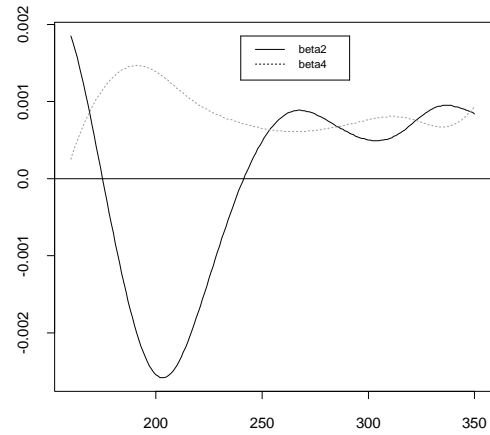


Figure 3: (a) Time series plot of Daily Hospital Admissions. (b) Scatterplot of Y_t vs Y_{t-1} . (c) Smoothed parametric estimation of β_1 . (d) Smoothed estimation of β_2 and β_4 .